

C O N T E N T S

Met	Lys	Ala	Leu	Arg	Ala	Val	Leu	Leu	Ile	Leu	Leu	Ser	Gly	Gln	
1				5					10				15		
Pro	Gly	Ser	Ser	Trp	Ala	Gln	Glu	Ala	Gly	Asp	Val	Asp	Leu	Glu	Leu
			20					25					30		
Glu	Arg	Tyr	Ser	Tyr	Asp	Asp	Asp	Gly	Asp	Asp	Asp	Asp	Asp	Asp	Asp
		35					40					45			
Glu	Glu	Glu	Glu	Glu	Glu	Glu	Thr	Asn	Met	Ile	Pro	Gly	Ser	Arg	Asp
	50					55					60				
Arg	Ala	Pro	Pro	Leu	Gln	Cys	Tyr	Phe	Cys	Gln	Val	Leu	His	Ser	Gly
65					70					75					80
Glu	Ser	Cys	Asn	Glu	Thr	Gln	Arg	Cys	Ser	Ser	Ser	Lys	Pro	Phe	Cys
			85						90					95	
Ile	Thr	Val	Ile	Ser	His	Gly	Lys	Thr	Asp	Thr	Gly	Val	Leu	Thr	Thr
			100					105					110		
Tyr	Ser	Met	Trp	Cys	Thr	Asp	Thr	Cys	Gln	Pro	Ile	Val	Lys	Thr	Val
		115					120					125			
Asp	Ser	Thr	Gln	Met	Thr	Gln	Thr	Cys	Cys	Gln	Ser	Thr	Leu	Cys	Asn
	130					135					140				
Ile	Pro	Pro	Trp	Gln	Ser	Pro	Gln	Ile	His	Asn	Pro	Leu	Gly	Gly	Arg
145					150					155					160
Ala	Asp	Ser	Pro	Leu	Lys	Gly	Gly	Thr	Arg	His	Pro	Gln	Gly	Asp	Arg
				165					170					175	
Phe	Ser	His	Pro	Gln	Val	Val	Lys	Val	Thr	His	Pro	Gln	Ser	Asp	Gly
			180					185					190		
Ala	His	Leu	Ser	Lys	Gly	Gly	Lys	Ala	Asn	Gln	Pro	Gln	Gly	Asn	Gly
		195					200					205			
Ala	Gly	Phe	Pro	Ala	Gly	Trp	Ser	Lys	Phe	Gly	Asn	Val	Val	Leu	Leu
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Leu	Thr	Phe	Leu	Thr	Ser	Leu	Trp	Ala	Ser	Gly	Ala				

235

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Ala Val Leu Leu Ile Leu Leu Leu Ser Gly Gln Pro Gly Ser Ser Trp
10 15 20

gca caa gaa gct ggc gat gtg gac ctg gag cta gag cgc tac agc tac 152
Ala Gln Glu Ala Gly Asp Val Asp Leu Glu Leu Glu Arg Tyr Ser Tyr
25 30 35

gat gat gac ggt gat gac gat gat gac gat gat gaa gaa gag gaa gag 200
Asp Asp Asp Gly Asp Asp Asp Asp Asp Asp Asp Glu Glu Glu Glu Glu
40 45 50

gag gag acc aac atg atc cct ggc agc agg gac aga gca ccg cct cta 248
Glu Glu Thr Asn Met Ile Pro Gly Ser Arg Asp Arg Ala Pro Pro Leu
55 60 65

cag tgc tac ttc tgc caa gtg ctt cac agc ggg gag agc tgc aac gag 296
Gln Cys Tyr Phe Cys Gln Val Leu His Ser Gly Glu Ser Cys Asn Glu
70 75 80 85

106⁵ ⁵aca⁵ cag aga tgc tcc agc agc aag ccc⁵ ttc tgt atc aca gtc atc tcc 344
Thr Gln Arg Cys Ser Ser Ser Lys Pro Phe Cys Ile Thr Val Ile Ser
90 95 100

cat ggc aaa act gac aca ggt gtc ctg acg acc tac tcc atg tgg tgt 392
His Gly Lys Thr Asp Thr Gly Val Leu Thr Thr Tyr Ser Met Trp Cys
105 110 115

act gat acc tgc cag ccc atc gtg aag aca gtg gac agc acc caa atg 440
Thr Asp Thr Cys Gln Pro Ile Val Lys Thr Val Asp Ser Thr Gln Met
120 125 130

acc cag acc tgt tgc cag tcc aca ctc tgc aat att cca ccc tgg cag 488
Thr Gln Thr Cys Cys Gln Ser Thr Leu Cys Asn Ile Pro Pro Trp Gln
135 140 145

gagc ccc caa atc cac aac cct ctg ggt ggc cgg gca gac agc ccc ttg 536
 Ser Pro Gln Ile His Asn Pro Leu Gly Gly Arg Ala Asp Ser Pro Leu
 150 155 160 165

aag ggt ggg acc aga cat cct caa ggt gac agg ttt agc cac ccc cag 584
Lys Gly Gly Thr Arg His Pro Gln Gly Asp Arg Phe Ser His Pro Gln
170 175 180

ggt ggc aag gct aac cag ccc cag gga aat ggg gcc gga ttc cct gca 680
 Gly Gly Lys Ala Asn Gln Pro Gln Gly Asn Gly Ala Gly Phe Pro Ala
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<212> DNA
<213> Artificial Sequence
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Questions **A**nswers **C**onclusions **E**xercises **P**roblems **S**tudies **T**estimonials **V**ideos **W**orkbooks

18

25

25

22

21

<210> 15
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<212> DNA
<213> Artificial Sequence

<220>
<223> synthetic

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gtccacagga	ctgctggatg	aggtacacca	catgactatg	gacggtcggg	tagcacttct	420
gtcacctgtc	gtggggttac	tgggtctgga	caacggtcag	gtgtgagacg	ttataagggtg	480
ggaccgtctc	gggggtttag	gtggtggggag	accacccggc	ccgtctgtcg	gggaacttcc	540
caccctgggtc	tgtaggagt	ccactgtcca	aatcggtggg	ggtccaacag	ttccaatgag	600
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